

SEQUENCE LISTING Max-Planck-Gesellschaft zur Förderung der Wissenschaften <110> Cell line comprising vector encoding truncated FLK-1 receptor <120> Div. of US 09/766 678 (Pat. No. 6 872 699) 44995P US <130> US 10/799,782 <140> . <141> 2009-06-05 <160> PatentIn version 3.3 <170> <210> <211> 5470 <212> DNA <213> Unknown <220> <223> SEQ ID NO: 1 <220> <221> CDS <222> (286) . . (4386) <400> tatagggcga attgggtacg ggacccccct cgaggtcgac ggtatcgata agcttgatat 60 cgaattcggg cccagactgt gtcccgcagc cgggataacc tggctgaccc gattccgcgg 120 acaccgctga cagccgcgc tggagccagg gcgccggtgc cccgcgctct ccccggtctt 180 240 gcgctgcggg ggccataccg cctctgtgac ttctttgcgg gccagggacg gagaaggagt 297 ctqtqcctqa gaaactgggc tctgtgccca ggcgcgaggt gcagg atg gag agc aag Met Glu Ser Lys gcg ctg cta gct gtc gct ctg tgg ttc tgc gtg gag acc cga gcc gcc 345 Ala Leu Leu Ala Val Ala Leu Trp Phe Cys Val Glu Thr Arg Ala Ala tct gtg ggt ttg act ggc gat ttt ctc cat ccc ccc aag ctc agc aca 393 Ser Val Gly Leu Thr Gly Asp Phe Leu His Pro Pro Lys Leu Ser Thr cag aaa gac ata ctg aca att ttg gca aat aca acc ctt cag att act 441 Gln Lys Asp Ile Leu Thr Ile Leu Ala Asn Thr Thr Leu Gln Ile Thr 489 tgc agg gga cag cgg gac ctg gac tgg ctt tgg ccc aat gct cag cgt Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro Asn Ala Gln Arg 60 gat tot gag gaa agg gta ttg gtg act gaa tgc ggc ggt ggt gac agt 537 Asp Ser Glu Glu Arg Val Leu Val Thr Glu Cys Gly Gly Asp Ser atc ttc tgc aaa aca ctc acc att ccc agg gtg gtt gga aat gat act 585 Ile Phe Cys Lys Thr Leu Thr Ile Pro Arg Val Val Gly Asn Asp Thr 95 100 90 85 gga gcc tac aag tgc tcg tac cgg gac gtc gac ata gcc tcc act gtt 633 Gly Ala Tyr Lys Cys Ser Tyr Arg Asp Val Asp Ile Ala Ser Thr Val

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Asn Ala Gln Arg Asp Ser Glu Glu Arg Val Leu Val Thr Glu Cys Gly

Gly Gly Asp Ser Ile Phe Cys Lys Thr Leu Thr Ile Pro Arg Val Val

Gly Asn Asp Thr Gly Ala Tyr Lys Cys Ser Tyr Arg Asp Val Asp Ile

Ala Ser Thr Val Tyr Val Tyr Val Arg Asp Tyr Arg Ser Pro Phe Ile

Ala Ser Val Ser Asp Gln His Gly Ile Val Tyr Ile Thr Glu Asn Lys 130

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455

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Seite 13

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Leu Thr Ile Met Glu Val Thr Glu Arg Asp Ala Gly Asn Tyr Thr Val 385 390 395 400

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Val Tyr Ala Asn Pro Pro Leu His His Ile Gln Trp Tyr Trp Gln Leu 450 455 460

Glu Glu Ala Cys Ser Tyr Arg Pro Gly Gln Thr Ser Pro Tyr Ala Cys 465 470 475 480

Lys Glu Trp Arg His Val Glu Asp Phe Gln Gly Gly Asn Lys Ile Glu 485 490 495

Val Thr Lys Asn Gln Tyr Ala Leu Ile Glu Gly Lys Asn Lys Thr Val 500 505 510

Ser Thr Leu Val Ile Gln Ala Ala Asn Val Ser Ala Leu Tyr Lys Cys 515 520 525

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Glu Gln Glu Ser Val Ser Leu Leu Cys Thr Ala Asp Arg Asn Thr Phe 565 570 575

Glu Asn Leu Thr Trp Tyr Lys Leu Gly Ser Gln Ala Thr Ser Val His 580 585 590

Met Gly Glu Ser Leu Thr Pro Val Cys Lys Asn Leu Asp Ala Leu Trp Lys Leu Asn Gly Thr Met Phe Ser Asn Ser Thr Asn Asp Ile Leu Ile 615

Val Ala Phe Gln Asn Ala Ser Leu Gln Asp Gln Gly Asp Tyr Val Cys

Ser Ala Gln Asp Lys Lys Thr Lys Lys Arg His Cys Leu Val Lys Gln

Leu Ile Ile Leu Glu Arg Met Ala Pro Met Ile Thr Gly Asn Leu Glu 665

Asn Gln Thr Thr Thr Ile Gly Glu Thr Ile Glu Val Thr Cys Pro Ala

Ser Gly Asn Pro Thr Pro His Ile Thr Trp Phe Lys Asp Asn Glu Thr 700

Leu Val Glu Asp Ser Gly Ile Val Leu Arg Asp Gly Asn Arg Asn Leu

Thr Ile Arg Arg Val Arg Lys Glu Asp Gly Gly Leu Tyr Thr Cys Gln

Ala Cys Asn Val Leu Gly Cys Ala Arg Ala Glu Thr Leu Phe Ile Ile

Glu Gly Ala Gln Glu Lys Thr Asn Leu Glu Val Ile Ile Leu Val Gly

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Leu Gln Ile Thr Cys Arg Gly Gln Arg Asp Leu Asp Trp Leu Trp Pro

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Ala	Ser 130	Val	Ser	Asp	Gln	His 135	Gly	Ile	Val	Tyr	Ile 140	Thr	Glu	Asn	Lys
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Val	Ser	Leu	Cys	Ala 165	Arg	Tyr	Pro	Glu	Lys 170	Arg	Phe	Val	Pro	Asp 175	Gly
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Lys	Ile	Val 275	Asn	Arg	Asp	Val	Lys 280	Pro	Phe	Pro	Gly	Thr 285	Val	Ala	Lys
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Leu 385	Thr	Ile	Met	Glu	Val 390	Thr	Glu	Arg	Asp	Ala 395	Gly	Asn	Tyr	Thr	Val 400
Ile	Leu	Thr	Asn	Pro 405	Ile	Ser	Met	Glu	Lys 410	Gln	Ser	His	Met	Val 415	Ser
Leu	Val	Val	Asn 420	Val	Pro	Pro	Gln	Ile 425	Gly	Glu	Lys	Ala	Leu 430	Ile	Ser
Pro	Met	Asp 435	Ser	Tyr	Gln	Tyr	Gly 440	Thr	Met	Gln	Thr	Leu 445	Thr	Cys	Thr
Val	туг 450	Ala	Asn	Pro	Pro	Leu 455	His	His	Ile	Gln	Trp 460	Tyr	Trp	Gln	Leu
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Lys	Glu	Trp	Arg	His 485	Val	Glu	Asp	Phe	Gln 490	Gly	Gly	Asn	Lys	Ile 495	Glu
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Ser	Thr	Leu 515	Val	Ile	Gln	Ala	Ala 520	Asn	Val	Ser	Ala	Leu 525	Tyr	Lys	Cys
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Glu	Gln	Glu	Ser	Val 565	Ser	Leu	Leu	Cys	Thr 570	Ala	Asp	Arg	Asn	Thr 575	Phe
Glu	Asn	Leu	Thr 580	Trp	Tyr	Lys	Leu	Gly 585	Ser	Gln	Ala	Thr	Ser 590	Val	His
Met	Gly	Glu 595	Ser	Leu	Thr	Pro	Val 600	Cys	Lys	Asn	Leu	Asp 605	Ala	Leu	Trp
Lys	Leu 610	Asn	Gly	Thr	Met	Phe 615	Ser	Asn	Ser	Thr	Asn 620	Asp	Ile	Leu	Ile
Val 625	Ala	Phe	Gln	Asn	Ala 630	Ser	Leu	Gln	Asp	Gln 635	Gly	Asp	Tyr	Val	Cys 640

Ser	Ala	GIn	Asp	Lys 645	Lys	rnr	rys	гÀг	650	nıs	Cys	Leu	vai	655	GIII
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Asn	Gln	Thr 675	Thr	Thr	Ile	Gly	Glu 680	Thr	Ile	Glu	Val	Thr 685	Cys	Pro	Ala
Ser	Gly 690	Asn	Pro	Thr	Pro	His 695	Ile	Thr	Trp	Phe	Lys 700	Asp	Asn	Glu	Thr
Leu 705	Val	Glu	Asp	Ser	Gly 710	Ile	Val	Leu	Arg	Asp 715	Gly	Asn	Arg	Asn	Leu 720
Thr	Ile	Arg	Arg	Val 725	Arg	Lys	Glu	Asp	Gly 730	Gly	Leu	Tyr	Thr	Cys 735	Gln
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Glu	Gly	Ala 755		Glu	Lys	Thr	Asn 760	Leu	Glu	Val	Ile	Ile 765	Leu	Val	Gly
Thr	Ala 770	Val	Ile	Ala	Met	Phe 775	Phe	Trp	Leu	Leu	Leu 780	Val	Ile	Val	Leu
Arg 785		Val	Lys	Arg	Ala 790	Asn	Glu	Gly	Glu	Leu 795	Lys	Thr	Gly	Tyr	Leu 800
Ser	Ile	Val	Met	Asp 805		Asp	Glu	Leu	Pro 810	Leu	Asp	Glu	Arg	Cys 815	Glu
Arg	Leu	Pro	Tyr 820		Ala	Ser	Lys	Trp 825	Glu	Phe	Pro	Arg	Asp 830	Arg	Leu
Lys	Leu	Gly 835		Pro	Leu	Gly	Arg 840	Gly	Ala	Phe	Gly	Gln 845	Val	Ile	Glu
Ala	Asp 850		Phe	Gly	· Ile	Asp 855	Lys	Thr	Ala	Thr	Cys 860	Lys	Thr	Val	Ala
Val 865	_	. Met	. Leu	Lys	61u 870		⁄ Ala	Thr	His	Ser 875	Glu	His	Arg	Ala	Leu 880
Met	: Ser	- Glu	ı Lev	Lys 885		e Lev	ı Ile	His	890	Gly	His	His	Leu	Asn 895	Val
Va]	Asn	ı Lei	1 Let 900		Ala	Суз	Thr	Lys 905		Gly	gly	Pro	Leu 910	Met	Val

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- Lys Arg Asn Glu Phe Val Pro Tyr Lys Ser Lys Gly Ala Arg Phe Arg 930 935 940
- Gln Gly Lys Asp Tyr Val Gly Glu Leu Ser Val Asp Leu Lys Arg Arg 945 950 955 960
- Leu Asp Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly Phe Val 965 970 975
- Glu Glu Lys Ser Leu Ser Asp Val Glu Glu Glu Glu Ala Ser Glu Glu 980 985 990
- Leu Tyr Lys Asp Phe Leu Thr Leu Glu His Leu Ile Cys Tyr Ser Phe 995 1000 1005
- Gln Val Ala Lys Gly Met Glu Phe Leu Ala Ser Arg Lys Cys Ile 1010 1015 1020
- His Arg Asp Leu Ala Ala Arg Asn Ile Leu Leu Ser Glu Lys Asn 1025 1030 1035
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- Trp Met Ala Pro Glu Thr Ile Phe Asp Arg Val Tyr Thr Ile Gln 1070 1080
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- Cys Arg Arg Leu Lys Glu Gly Thr Arg Met Arg Ala Pro Asp Tyr 1115 1120 1125
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- Asp Pro Asn Gln Arg Pro Ser Phe Ser Glu Leu Val Glu His Leu 1145 1150 1155
- Gly Asn Leu Leu Gln Ala Asn Ala Gln Gln Asp Gly Lys Asp Tyr 1160 1165 1170
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1175 1180 1185

Gly Leu Ser Leu Pro Thr Ser Pro Val Ser Cys Met Glu Glu 1190 1195 1200

Glu Val Cys Asp Pro Lys Phe His Tyr Asp Asn Thr Ala Gly Ile 1205 1210 1215

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Val Lys Thr Phe Glu Asp Ile Pro Leu Glu Glu Pro Glu Val Lys 1240

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Ser Glu Glu Leu Lys Thr Leu Glu Asp Arg Asn Lys Leu Ser Pro 1270

Ser Phe Gly Gly Met Met Pro Ser Lys Ser Arg Glu Ser Val Ala 1285

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Lys Phe Gly Asn Leu Ser Thr Tyr Leu Arg Gly Lys Arg Asn Glu Phe

Val Pro Tyr Lys Ser Lys Gly Ala Arg Phe Arg Gln Gly Lys Asp Tyr Val Gly Glu Leu Ser Val Asp Leu Lys Arg Arg Leu Asp Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly Phe Val Glu Glu Lys Ser Leu Ser Asp Val Glu Glu Glu Ala Ser Glu Glu Leu Tyr Lys Asp Phe Leu Thr Leu Glu His Leu Ile Cys Tyr Ser Phe Gln Val Ala Lys Gly 120 Met Glu Phe Leu Ala Ser Arg Lys Cys Ile His Arg Asp Leu Ala Ala Arg Asn Ile Leu Leu Ser Glu Lys Asn Val Val Lys Ile Cys Asp Phe Gly Leu Ala Arg Asp Ile Tyr Lys Asp Pro Asp Tyr Val Arg Lys Gly
165 170 175 Asp Ala Arg Leu <210> 10 <211> 180 <212> PRT <213> Unknown <220> <223> SEQ ID NO: 10 Ile Leu Ile His Ile Gly His His Leu Asn Val Val Asn Leu Leu Gly Ala Cys Thr Lys Pro Gly Gly Pro Leu Met Val Ile Val Glu Phe Cys Lys Phe Asp Asn Leu Ser Thr Tyr Leu Arg Ser Lys Arg Asn Glu Phe Val Pro Tyr Lys Thr Lys Gly Ala Arg Phe Arg Gln Gly Lys Asp Tyr Val Gly Ala Ile Pro Val Asp Leu Lys Arg Arg Leu Asp Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly Phe Val Glu Glu Lys Ser Leu 90

Ser Asp Val Glu Glu Glu Ala Pro Glu Asp Leu Tyr Lys Asp Phe 105 100 Leu Thr Leu Glu His Leu Ile Cys Tyr Ser Phe Gln Val Ala Lys Gly Met Glu Phe Leu Ala Ser Arg Lys Cys Ile His Arg Asp Leu Ala Ala Arg Asn Ile Leu Leu Ser Glu Lys Asn Val Val Lys Ile Cys Asp Phe Gly Leu Ala Arg Asp Ile Tyr Lys Asp Pro Asp Tyr Val Arg Lys Gly Asp Ala Arg Leu <210> 11 <211> 180 <212> PRT <213> Unknown <220> <223> SEQ ID NO: 11 <400> 11 Ile Leu Ile His Ile Gly His His Leu Asn Val Val Asn Leu Leu Gly Ala Cys Thr Lys Pro Gly Gly Pro Leu Met Val Ile Val Glu Phe Cys Lys Phe Gly Asn Leu Ser Thr Tyr Leu Arg Gly Lys Arg Asn Glu Phe Val Pro Tyr Lys Ser Lys Gly Ala Arg Phe Arg Ser Gly Lys Asp Tyr Val Gly Glu Leu Ser Val Asp Leu Lys Arg Arg Leu Asp Ser Ile Thr Ser Ser Gln Ser Ser Ala Ser Ser Gly Phe Val Glu Glu Lys Ser Leu Ser Asp Val Glu Glu Glu Glu Ala Ser Glu Glu Leu Tyr Lys Asp Phe Leu Thr Leu Glu His Leu Ile Cys Tyr Ser Phe Gln Val Ala Lys Gly

Seite 29

Met Glu Phe Leu Ala Ser Arg Lys Cys Ile His Arg Asp Leu Ala Ala

135

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Gly Leu Ala Arg Asp Ile Tyr Lys Asp Pro Asp Tyr Val Arg Lys Gly 165 170 175

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